

0570  
0528

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# 10

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/940,921B

DATE: 06/03/2002

TIME: 13:52:47

Input Set : A:\LEX-0227-USA SEQLIST.txt

Output Set: N:\CRF3\06032002\I940921B.raw

4 <110> APPLICANT: Friddle, Carl Johan  
5 Hilbun, Erin  
6 Nepomnichy, Boris  
7 Hu, Yi

9 <120> TITLE OF INVENTION: Novel Human Kinase Proteins and Polynucleotides Encoding the

Same

11 <130> FILE REFERENCE: LEX-0227-USA  
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/940,921B  
C--> 13 <141> CURRENT FILING DATE: 2002-05-21  
13 <150> PRIOR APPLICATION NUMBER: US 60/229,280  
14 <151> PRIOR FILING DATE: 2000-08-31  
16 <160> NUMBER OF SEQ ID NOS: 10  
18 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
20 <210> SEQ ID NO: 1  
21 <211> LENGTH: 2052  
22 <212> TYPE: DNA  
23 <213> ORGANISM: homo sapiens  
25 <400> SEQUENCE: 1  
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27 gctaaaggaa aatcagatag caagcactgt gtcataaaag agatcaattt tgaaaagatg 120  
28 cccataacaag aaaaagaagc ttcaaaagaaa gaagtgattc ttctggaaaa gatgaaacat 180  
29 cccaaacattt tagccttcaattt caagagaatg gcaggctgtt tattttaatg 240  
30 gaatattgtt atggagggaa tctcatgaaa aggatcaata gacaacgggg tttgttattt 300  
31 agtgaagatc agatccctgg ttggttttaa cagatttctc taggactaaa acatattcat 360  
32 gacaggaaga tattacacag ggacataaaa gctcagaaca tttttcttag caagaacgga 420  
33 atggtggcaa agcttggggaa ctttggata gcaagagtcc tgaataattt catggaaactt 480  
34 gctcgaactt gtatttggaa accttactac ctgtccccag agatctgtca gaataaaaccc 540  
35 tacaacaata aaacggatata ttggctcttt ggctgtgtct tatatgagct ctgcacactt 600  
36 aaacatcctt ttgagggtaa caacttacag cagctggttc tgaagattt tcaagcacat 660  
37 tttggcccaa tatctccggg gttttctcggtt gagctccatt ctttgatatt tcagctctt 720  
38 caagtatctc ctgcagaccg accatccata aattccattt tgaaaaggcc ttttttagag 780  
39 aatcttattc ccaaataattt gactcctgag gtcattcagg aagaattcag tcacatgctt 840  
40 atatgcagag caggagcc agcttctcgat catgtggaa aggtggtcca gaagtgtaaa 900  
41 atacaaaaag tgagattcca gggaaagtgc ccaccaagat caaggatattc tttgtccatt 960  
42 aaaaggaatg ctatattgca tagaaatgaa tggagaccac cagctggagc ccagaaggcc 1020  
43 agatctataa aaatgtataga aagacccaaa attgtctgtc tctgtggaca ttatgtattt 1080  
44 tattatgctc aacttgcata gctgaggagg agagccaca aaccaagtta tcaccctatt 1140  
45 cctcaagaaa atactggatc tgaggattac ggtcaggaaa cgaggcatgg tccatccccca 1200  
46 agtcaatggc ctgctgatc cttcagaga aaattttaa gtcacaaata taagtgtaaa 1260  
47 gtggagaagc aattgggtct tcgtccatct tctgcccggc caaattacaa ccagagacaa 1320  
48 gagctaagaa gtaatggaga agagcctaga ttccaggagc tgccatattt gaaaaacggaa 1380  
49 atgaaggaac aggaatattt gaagcgttta gaggaaatac gccaacagta ccacaatgac 1440  
50 atgaaagaaa tttagaaagaa gatggggaga gaaccagagg agaactcaaa aataagtcat 1500  
51 aaaacctatt tggtaagaa gatgtccatc aagatgcattc tgagggagaa 1560

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52	gcacctgtgc	aggacattga	aaaagacttg	aaacaaatga	ggcttcagaa	cacaaaggaa	1620										
53	agtaaaaatc	cagaacagaa	atataaagct	aagaaggggg	taaaatttga	aattaattta	1680										
54	gacaaatgt	tttctgtat	aaacatcctc	caagaggaag	aggcaatgga	tataccaaat	1740										
55	gaaacttga	ccttgagga	tggcatgaag	tttaaggaat	atgaatgtgt	aaaggagcat	1800										
56	ggagattata	cagacaaaagc	atttgaaaaa	cttcactgccc	cagaagcagg	gttttccacg	1860										
57	cagactgtag	ctgctgtggg	aaacaggagg	cagtggatg	gaggagcggcc	tcagactctg	1920										
58	ctgcagatga	tggcagtggc	cgacatcacc	tccactgccc	ccacggggcc	tgacagttag	1980										
59	tctgtgctta	gcgtcagtcg	tcaggaaggg	aagaccaagg	accctacag	cccagtgctc	2040										
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63	<211> LENGTH: 683																
64	<212> TYPE: PRT																
65	<213> ORGANISM: homo sapiens																
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69	1					5				10					15		
70	Lys	Ala	Tyr	Leu	Ala	Lys	Gly	Lys	Ser	Asp	Ser	Lys	His	Cys	Val	Ile	
71						20				25					30		
72	Lys	Glu	Ile	Asn	Phe	Glu	Lys	Met	Pro	Ile	Gln	Glu	Lys	Glu	Ala	Ser	
73						35				40					45		
74	Lys	Lys	Glu	Val	Ile	Leu	Leu	Glu	Lys	Met	Lys	His	Pro	Asn	Ile	Val	
75						50				55					60		
76	Ala	Phe	Phe	Asn	Ser	Phe	Gln	Glu	Asn	Gly	Arg	Leu	Phe	Ile	Val	Met	
77						65				70					75		80
78	Glu	Tyr	Cys	Asp	Gly	Gly	Asp	Leu	Met	Lys	Arg	Ile	Asn	Arg	Gln	Arg	
79						85				90					95		
80	Gly	Val	Leu	Phe	Ser	Glu	Asp	Gln	Ile	Leu	Gly	Trp	Phe	Val	Gln	Ile	
81						100				105					110		
82	Ser	Leu	Gly	Leu	Lys	His	Ile	His	Asp	Arg	Lys	Ile	Leu	His	Arg	Asp	
83						115				120					125		
84	Ile	Lys	Ala	Gln	Asn	Ile	Phe	Leu	Ser	Lys	Asn	Gly	Met	Val	Ala	Lys	
85						130				135					140		
86	Leu	Gly	Asp	Phe	Gly	Ile	Ala	Arg	Val	Leu	Asn	Asn	Ser	Met	Glu	Leu	
87						145				150					155		160
88	Ala	Arg	Thr	Cys	Ile	Gly	Thr	Pro	Tyr	Tyr	Leu	Ser	Pro	Glu	Ile	Cys	
89						165				170					175		
90	Gln	Asn	Lys	Pro	Tyr	Asn	Asn	Lys	Thr	Asp	Ile	Trp	Ser	Leu	Gly	Cys	
91						180				185					190		
92	Val	Leu	Tyr	Glu	Leu	Cys	Thr	Leu	Lys	His	Pro	Phe	Glu	Gly	Asn	Asn	
93						195				200					205		
94	Leu	Gln	Gln	Leu	Val	Leu	Lys	Ile	Cys	Gln	Ala	His	Phe	Ala	Pro	Ile	
95						210				215					220		
96	Ser	Pro	Gly	Phe	Ser	Arg	Glu	Leu	His	Ser	Leu	Ile	Ser	Gln	Leu	Phe	
97						225				230					235		240
98	Gln	Val	Ser	Pro	Arg	Asp	Arg	Pro	Ser	Ile	Asn	Ser	Ile	Leu	Lys	Arg	
99						245				250					255		
100	Pro	Phe	Leu	Glu	Asn	Leu	Ile	Pro	Lys	Tyr	Leu	Thr	Pro	Glu	Val	Ile	
101						260				265					270		
102	Gln	Glu	Glu	Phe	Ser	His	Met	Leu	Ile	Cys	Arg	Ala	Gly	Ala	Pro	Ala	

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Input Set : A:\LEX-0227-USA SEQLIST.txt  
Output Set: N:\CRF3\06032002\I940921B.raw

103	275	280	285	
104	Ser Arg His Ala Gly Lys Val Val Gln Lys Cys Lys Ile Gln Lys Val			
105	290	295	300	
106	Arg Phe Gln Gly Lys Cys Pro Pro Arg Ser Arg Ile Ser Val Pro Ile			
107	305	310	315	320
108	Lys Arg Asn Ala Ile Leu His Arg Asn Glu Trp Arg Pro Pro Ala Gly			
109	325	330	335	
110	Ala Gln Lys Ala Arg Ser Ile Lys Met Ile Glu Arg Pro Lys Ile Ala			
111	340	345	350	
112	Ala Val Cys Gly His Tyr Asp Tyr Tyr Tyr Ala Gln Leu Asp Met Leu			
113	355	360	365	
114	Arg Arg Arg Ala His Lys Pro Ser Tyr His Pro Ile Pro Gln Glu Asn			
115	370	375	380	
116	Thr Gly Val Glu Asp Tyr Gly Gln Glu Thr Arg His Gly Pro Ser Pro			
117	385	390	395	400
118	Ser Gln Trp Pro Ala Glu Tyr Leu Gln Arg Lys Phe Glu Ala Gln Gln			
119	405	410	415	
120	Tyr Lys Leu Lys Val Glu Lys Gln Leu Gly Leu Arg Pro Ser Ser Ala			
121	420	425	430	
122	Glu Pro Asn Tyr Asn Gln Arg Gln Glu Leu Arg Ser Asn Gly Glu Glu			
123	435	440	445	
124	Pro Arg Phe Gln Glu Leu Pro Phe Arg Lys Asn Glu Met Lys Glu Gln			
125	450	455	460	
126	Glu Tyr Trp Lys Gln Leu Glu Glu Ile Arg Gln Gln Tyr His Asn Asp			
127	465	470	475	480
128	Met Lys Glu Ile Arg Lys Lys Met Gly Arg Glu Pro Glu Glu Asn Ser			
129	485	490	495	
130	Lys Ile Ser His Lys Thr Tyr Leu Val Lys Lys Ser Asn Leu Pro Val			
131	500	505	510	
132	His Gln Asp Ala Ser Glu Gly Glu Ala Pro Val Gln Asp Ile Glu Lys			
133	515	520	525	
134	Asp Leu Lys Gln Met Arg Leu Gln Asn Thr Lys Glu Ser Lys Asn Pro			
135	530	535	540	
136	Glu Gln Lys Tyr Lys Ala Lys Lys Gly Val Lys Phe Glu Ile Asn Leu			
137	545	550	555	560
138	Asp Lys Cys Ile Ser Asp Glu Asn Ile Leu Gln Glu Glu Ala Met			
139	565	570	575	
140	Asp Ile Pro Asn Glu Thr Leu Thr Phe Glu Asp Gly Met Lys Phe Lys			
141	580	585	590	
142	Glu Tyr Glu Cys Val Lys Glu His Gly Asp Tyr Thr Asp Lys Ala Phe			
143	595	600	605	
144	Glu Lys Leu His Cys Pro Glu Ala Gly Phe Ser Thr Gln Thr Val Ala			
145	610	615	620	
146	Ala Val Gly Asn Arg Arg Gln Trp Asp Gly Gly Ala Pro Gln Thr Leu			
147	625	630	635	640
148	Leu Gln Met Met Ala Val Ala Asp Ile Thr Ser Thr Cys Pro Thr Gly			
149	645	650	655	
150	Pro Asp Ser Glu Ser Val Leu Ser Val Ser Arg Gln Glu Gly Lys Thr			
151	660	665	670	

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/940,921B

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Input Set : A:\LEX-0227-USA SEQLIST.txt  
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152 Lys Asp Pro Tyr Ser Pro Val Leu Ile Leu Met  
 153 675 680  
 155 <210> SEQ ID NO: 3  
 156 <211> LENGTH: 1965  
 157 <212> TYPE: DNA  
 158 <213> ORGANISM: homo sapiens  
 160 <400> SEQUENCE: 3

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162	gctaaaggga	aatcagatag	caagcactgt	gtcataaaag	agatcaattt	tgaaaagatg	120										
163	cccatacaag	aaaaagaagc	ttcaaagaaa	gaagtgattc	ttctggaaaa	gatgaaacat	180										
164	cccaacattt	tagccttctt	caattcattt	caagagaatg	gcaggctgtt	tattttaatg	240										
165	aatatttgtt	atggagggga	tctcatgaaa	aggatcaata	gacaacgggg	tgtgttattt	300										
166	agtgaagatc	agatcctcg	ttggtttgtt	cagatttctc	taggactaaa	acatattcat	360										
167	gacaggaaga	tattacacag	ggacataaaaa	gctcagaaca	tttttcttag	caagaacgga	420										
168	atggtggcaa	agcttgggaa	cttttgtata	gcaagagtcc	tgaataattt	catggaactt	480										
169	gtcgaactt	gtatttggAAC	accttactac	ctgtcccccag	agatctgtca	gaataaaccc	540										
170	tacaacaata	aaacggatata	ttggtcttctt	ggctgtgtct	tatatgagct	ctgcacactt	600										
171	aaacatcctt	ttgagggtaa	caacttacag	cagctggttc	tgaagattt	tcaagcacat	660										
172	tttgccccaa	tatctccggg	gttttctcg	gagctccatt	ctttgatata	tcaagtcttt	720										
173	caagtatctc	ctcgagaccg	accatccata	aattccattt	tgaaaaggcc	cttttagag	780										
174	aatcttattt	ccaaatattt	gactcctgag	gtcattcagg	aagaatttc	tcacatgctt	840										
175	atatgcagag	caggagcgc	agcttctcg	catgtggga	aggtggtcca	gaagtgtaaa	900										
176	ataaaaaaaag	ttagatttca	gggaaagtgc	ccaccaagat	caaggatata	tgtgccaatt	960										
177	aaaaggaatg	ctatatttgc	tagaaatgaa	tggagaccac	cagctggagc	ccagaaggcc	1020										
178	agatctataa	aatgtataga	aagacccaaa	attgtgtct	tctgtggaca	ttatgattat	1080										
179	tattatgctc	aacttggat	gctgaggagg	agagccaca	aaccaagttt	tcaccctatt	1140										
180	cctcaagaaa	atactggagt	tgaggattac	ggtcaggaaa	cgaggcatgg	tccatcccc	1200										
181	agtcaatggc	ctgctgagta	ccttcagaga	aaatttgaag	ctcaacaata	taagttgaaa	1260										
182	gtggagaagc	aattgggtct	tcgtccatct	tctggcggac	caaattacaa	ccagagacaa	1320										
183	gagctaagaa	gtaatggaga	agagcctaga	ttccaggagc	tgccatttt	gaaaaacgaa	1380										
184	atgaaggaac	aggagaactc	aaaaataagt	cataaaacct	attttgtgaa	gaagagtaac	1440										
185	ctgcctgtcc	atcaagatgc	atctgaggg	gaagcacctg	tgcaggacat	tgaaaaagac	1500										
186	ttgaaaacaaa	tgaggcttca	gaacacaaa	gaaagtaaaa	atccagaaca	gaaatataaa	1560										
187	gctaagaagg	gggtaaaatt	tgaaattat	ttagacaaaat	gtatttctgt	tgaaaacatc	1620										
188	ctccaagagg	aagaggcaat	ggatatacca	aatgaaaactt	tgacctttgt	ggatggcatg	1680										
189	aagtttaagg	aatatgtatg	tgtaaaggag	catggagatt	atacagacaa	agcatttgaa	1740										
190	aaacttcaact	gcccagaagc	agggtttcc	acgcagactg	tagctgtgt	ggaaacagg	1800										
191	aggcagtggg	atggaggagc	gcctcagact	ctgctgcaga	tgtggcagt	ggccgacatc	1860										
192	acctccaccc	gccccacggg	gcctgacagt	gagtcgtgc	ttagcgtcag	tcgtcaggaa	1920										
193	gggaagacca	aggacccgt	cagcccgat	ctcatcctga	tgtga		1965										
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196	<211>	LENGTH: 654															
197	<212>	TYPE: PRT															
198	<213>	ORGANISM: homo sapiens															
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202	1				5				10						15		
203	Lys	Ala	Tyr	Leu	Ala	Lys	Gly	Lys	Ser	Asp	Ser	Lys	His	Cys	Val	Ile	
204					20				25						30		

RAW SEQUENCE LISTING  
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Input Set : A:\LEX-0227-USA SEQLIST.txt  
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205 Lys Glu Ile Asn Phe Glu Lys Met Pro Ile Gln Glu Lys Glu Ala Ser  
206 35 40 45  
207 Lys Lys Glu Val Ile Leu Leu Glu Lys Met Lys His Pro Asn Ile Val  
208 50 55 60  
209 Ala Phe Phe Asn Ser Phe Gln Glu Asn Gly Arg Leu Phe Ile Val Met  
210 65 70 75 80  
211 Glu Tyr Cys Asp Gly Gly Asp Leu Met Lys Arg Ile Asn Arg Gln Arg  
212 85 90 95  
213 Gly Val Leu Phe Ser Glu Asp Gln Ile Leu Gly Trp Phe Val Gln Ile  
214 100 105 110  
215 Ser Leu Gly Leu Lys His Ile His Asp Arg Lys Ile Leu His Arg Asp  
216 115 120 125  
217 Ile Lys Ala Gln Asn Ile Phe Leu Ser Lys Asn Gly Met Val Ala Lys  
218 130 135 140  
219 Leu Gly Asp Phe Gly Ile Ala Arg Val Leu Asn Asn Ser Met Glu Leu  
220 145 150 155 160  
221 Ala Arg Thr Cys Ile Gly Thr Pro Tyr Tyr Leu Ser Pro Glu Ile Cys  
222 165 170 175  
223 Gln Asn Lys Pro Tyr Asn Asn Lys Thr Asp Ile Trp Ser Leu Gly Cys  
224 180 185 190  
225 Val Leu Tyr Glu Leu Cys Thr Leu Lys His Pro Phe Glu Gly Asn Asn  
226 195 200 205  
227 Leu Gln Gln Leu Val Leu Lys Ile Cys Gln Ala His Phe Ala Pro Ile  
228 210 215 220  
229 Ser Pro Gly Phe Ser Arg Glu Leu His Ser Leu Ile Ser Gln Leu Phe  
230 225 230 235 240  
231 Gln Val Ser Pro Arg Asp Arg Pro Ser Ile Asn Ser Ile Leu Lys Arg  
232 245 250 255  
233 Pro Phe Leu Glu Asn Leu Ile Pro Lys Tyr Leu Thr Pro Glu Val Ile  
234 260 265 270  
235 Gln Glu Glu Phe Ser His Met Leu Ile Cys Arg Ala Gly Ala Pro Ala  
236 275 280 285  
237 Ser Arg His Ala Gly Lys Val Val Gln Lys Cys Lys Ile Gln Lys Val  
238 290 295 300  
239 Arg Phe Gln Gly Lys Cys Pro Pro Arg Ser Arg Ile Ser Val Pro Ile  
240 305 310 315 320  
241 Lys Arg Asn Ala Ile Leu His Arg Asn Glu Trp Arg Pro Pro Ala Gly  
242 325 330 335  
243 Ala Gln Lys Ala Arg Ser Ile Lys Met Ile Glu Arg Pro Lys Ile Ala  
244 340 345 350  
245 Ala Val Cys Gly His Tyr Asp Tyr Tyr Tyr Ala Gln Leu Asp Met Leu  
246 355 360 365  
247 Arg Arg Arg Ala His Lys Pro Ser Tyr His Pro Ile Pro Gln Glu Asn  
248 370 375 380  
249 Thr Gly Val Glu Asp Tyr Gly Gln Glu Thr Arg His Gly Pro Ser Pro  
250 385 390 395 400  
251 Ser Gln Trp Pro Ala Glu Tyr Leu Gln Arg Lys Phe Glu Ala Gln Gln  
252 405 410 415  
253 Tyr Lys Leu Lys Val Glu Lys Gln Leu Gly Leu Arg Pro Ser Ser Ala

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/940,921B

DATE: 06/03/2002

TIME: 13:52:48

Input Set : A:\LEX-0227-USA SEQLIST.txt  
Output Set: N:\CRF3\06032002\I940921B.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date